

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: March 24, 2004, 22:33:46 ; Search time 45 Seconds
(without alignments)
1801.178 Million cell updates/sec

Title: US-09-900-038a-1
Perfect score: 1590
Sequence: 1 MNYSIMSYNNEPLNVRDS.....LINDINILVTLFGGEKQSD 313

Scoring table: BLOSUM62
Gapop 10.0, Gapext 0.5

Searched: 1049977 seqs, 258955339 residues

Total number of hits satisfying chosen parameters: 1049977

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :

1: /cgn2_6/ptodata/2/pubppaa/US07_PUBCOMB.pep.*
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11: /cgn2_6/ptodata/2/pubppaa/US09C_NEW_PUB.pep.*
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13: /cgn2_6/ptodata/2/pubppaa/US10_PUBCOMB.pep.*
14: /cgn2_6/ptodata/2/pubppaa/US10C_PUBCOMB.pep.*
15: /cgn2_6/ptodata/2/pubppaa/US10C_NEW_PUB.pep.*
16: /cgn2_6/ptodata/2/pubppaa/US60_NEW_PUB.pep.*
17: /cgn2_6/ptodata/2/pubppaa/US60_PUBCOMB.pep.*
18: /cgn2_6/ptodata/2/pubppaa/US60_PUBCOMB.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	1590	100.0	313	US-09-900-038a-1	Sequence 1, Appl1
2	367	23.1	333	US-10-282-122A-52072	Sequence 52072, A
3	348.5	23.9	336	US-10-282-122A-51835	Sequence 51835, A
4	317	19.9	333	US-10-282-122A-52169	Sequence 52169, A
5	240.5	15.1	297	US-09-738-626-3905	Sequence 3905, Ap
6	237	14.9	297	US-09-816-028A-31	Sequence 31, Appl
7	237	14.9	297	US-10-303-161-31	Sequence 31, Appl
8	237	14.9	297	US-10-303-118-31	Sequence 31, Appl
9	237	14.9	297	US-10-303-128-31	Sequence 31, Appl
10	237	14.9	297	US-10-303-134-31	Sequence 31, Appl
11	237	14.9	297	US-10-303-162-31	Sequence 31, Appl
12	234	14.7	301	US-09-816-028A-27	Sequence 27, Appl
13	234	14.7	301	US-10-303-161-27	Sequence 27, Appl
14	234	14.7	301	US-10-303-118-27	Sequence 27, Appl
15	234	14.7	301	US-10-303-128-27	Sequence 27, Appl

16	234	14.7	301	US-10-303-134-27	Sequence 27, Appl
17	234	14.7	301	US-10-303-162-27	Sequence 27, Appl
18	230.5	14.5	249	US-10-282-122A-53337	Sequence 53337, A
19	223	14.0	210	US-09-767-041-47	Sequence 47, Appl
20	218	13.7	874	US-10-282-122A-52215	Sequence 52215, A
21	216.5	13.6	1047	US-10-282-122A-56851	Sequence 56851, A
22	216	13.6	706	US-09-815-242-4950	Sequence 4950, Ap
23	216	13.6	713	US-10-282-122A-44526	Sequence 44526, A
24	216	13.6	713	US-10-282-122A-56852	Sequence 56852, A
25	216	13.6	715	US-09-815-242-10511	Sequence 10511, A
26	214	13.5	269	US-09-767-041-41	Sequence 41, Appl
27	210.5	13.2	348	US-10-654-528-3	Sequence 3, Appl
28	210.5	13.2	348	US-10-654-528-11	Sequence 11, Appl
29	210.5	13.2	348	US-10-007-267-3	Sequence 3, Appl
30	210.5	13.2	348	US-10-007-267-11	Sequence 11, Appl
31	210.5	13.2	348	US-10-096-129-3	Sequence 3, Appl1
32	210.5	13.2	348	US-10-096-129-8	Sequence 8, Appl1
33	210	13.2	322	US-09-767-041-34	Sequence 34, Appl
34	206.5	13.0	335	US-10-282-122A-74482	Sequence 74482, A
35	204.5	12.9	438	US-10-282-122A-60207	Sequence 60207, A
36	204	12.8	421	US-10-282-122A-48233	Sequence 48233, A
37	202.5	12.7	317	US-10-282-122A-73707	Sequence 73707, A
38	200	12.6	270	US-09-816-028A-39	Sequence 39, Appl
39	200	12.6	270	US-10-303-118-39	Sequence 39, Appl
40	200	12.6	270	US-10-303-128-39	Sequence 39, Appl
41	200	12.6	270	US-10-303-134-39	Sequence 39, Appl
42	200	12.6	270	US-10-303-162-39	Sequence 39, Appl
43	200	12.6	270	US-10-303-162-39	Sequence 39, Appl
44	200	12.6	337	US-10-282-122A-67193	Sequence 67193, A
45	196.5	12.4	294	US-10-282-122A-58582	Sequence 58582, A

ALIGNMENTS

RESULT 1
US-09-900-038a-1
Sequence 1, Application US/09900038A
Patent No. US20020142425A1
GENERAL INFORMATION:
APPLICANT: Miyake, Katsunide
APPLICANT: Watanabe, Masaki
APPLICANT: Iijima, Shinji
TITLE OF INVENTION: Beta 1,3-galactosyltransferase and DNA encoding the same
FILE REFERENCE: 766.53
CURRENT APPLICATION NUMBER: US/09/900,038A
PRIOR FILING DATE: 2001-09-21
PRIOR APPLICATION NUMBER: JP 2001-392
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1
LENGTH: 313
TYPE: PRT
ORGANISM: Streptococcus agalactiae Type 1b
US-09-900-038a-1

Query Match 100.0%; Score 1590; DB 9; Length 313;
Best Local Similarity 100.0%; Pred. No. 1,16-135;
Matches 313; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MNYSIMSYNNEPLNVRDSVSINQTLTDEFFITVDNPSRGDKOFLTEYSVVDRI 60
1 MNYSIMSYNNEPLNVRDSVSINQTLTDEFFITVDNPSRGDKOFLTEYSVVDRI 60
DB 1 MNYSIMSYNNEPLNVRDSVSINQTLTDEFFITVDNPSRGDKOFLTEYSVVDRI 60
QY 61 KIILNEENIGLASLNKAVKISKGEYIFRMDADISYPSRFDKQIRFMENSLDSATLI 120
61 KIILNEENIGLASLNKAVKISKGEYIFRMDADISYPSRFDKQIRFMENSLDSATLI 120
DB 61 KIILNEENIGLASLNKAVKISKGEYIFRMDADISYPSRFDKQIRFMENSLDSATLI 120
QY 121 ELIDQGNLVYKQREBNKTYLINDIRKMLNSIILAHPTWCYKKVFDLWGYRDLVPVE 180
121 ELIDQGNLVYKQREBNKTYLINDIRKMLNSIILAHPTWCYKKVFDLWGYRDLVPVE 180
DB 121 ELIDQGNLVYKQREBNKTYLINDIRKMLNSIILAHPTWCYKKVFDLWGYRDLVPVE 180

QY 181 DVDFATGALADPKIGLGNKVLQYRLNENG-SQTNKFOQYISALIQDFYKESYIDIT 240
DB 181 DVDFATGALADPKIGLGNKVLQYRLNENG-SQTNKFOQYISALIQDFYKESYIDIT 240
QY 241 KITNYPQEVYIKKRYQOELSKYFELKSTPSTIRKLYTCLVLYPKSPVLRLLINDINI 300
DB 241 KITNYPQEVYIKKRYQOELSKYFELKSTPSTIRKLYTCLVLYPKSPVLRLLINDINI 300
QY 301 LVKLFGKQKSD 313
DB 301 LVKLFGKQKSD 313

RESULT 2
US-10-282-122A-52072

/ Sequence 52072, Application US/10282122A
/ Publication No. US20040029129A1
/ GENERAL INFORMATION:
/ APPLICANT: Wang, Liangsu
/ APPLICANT: Zamudio, Carlos
/ APPLICANT: Malone, Cheryl
/ APPLICANT: Haselbeck, Robert
/ APPLICANT: Ohlsen, Kari
/ APPLICANT: Zyskind, Judith
/ APPLICANT: Wall, Daniel
/ APPLICANT: Trawick, John
/ APPLICANT: Carr, Grant
/ APPLICANT: Yamamoto, Robert
/ APPLICANT: Forsyth, R.
/ APPLICANT: Xu, H.
/ TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
/ FILE REFERENCE: EUTRA.034A
/ CURRENT APPLICATION NUMBER: US/10/282,122A
/ PRIOR FILING DATE: 2003-02-20
/ PRIOR APPLICATION NUMBER: 60/191,078
/ PRIOR FILING DATE: 2000-03-21
/ PRIOR APPLICATION NUMBER: 60/206,848
/ PRIOR FILING DATE: 2000-05-23
/ PRIOR APPLICATION NUMBER: 60/207,727
/ PRIOR FILING DATE: 2000-05-26
/ PRIOR APPLICATION NUMBER: 60/230,335
/ PRIOR FILING DATE: 2000-09-06
/ PRIOR APPLICATION NUMBER: 60/230,347
/ PRIOR FILING DATE: 2000-09-09
/ PRIOR APPLICATION NUMBER: 60/242,578
/ PRIOR FILING DATE: 2000-10-23
/ PRIOR APPLICATION NUMBER: 60/253,625
/ PRIOR FILING DATE: 2000-11-27
/ PRIOR APPLICATION NUMBER: 60/257,931
/ PRIOR FILING DATE: 2000-12-22
/ PRIOR APPLICATION NUMBER: 60/267,636
/ PRIOR FILING DATE: 2001-02-09
/ PRIOR APPLICATION NUMBER: 60/269,308
/ PRIOR FILING DATE: 2001-02-16
/ Remaining Prior Application data removed - See File Wrapper or PALM.
/ NUMBER OF SEQ ID NOS: 78614
/ SOFTWARE: PatentIn version 3.1
/ SEQ ID NO 52072
/ LENGTH: 333
/ TYPE: PRT
/ ORGANISM: Clostridium acetobutylicum
US-10-282-122A-52072

Query Match 23.1%, Score 367, DB 12, Length 333;
Best Local Similarity 33.3%, Pred. No. 6, Se-25;
Matches 93, Conservative 62, Mismatches 92, Indels 32, Gaps 10;
QY 4 STMSVNEPLNVRDSESTINOTLDPFEFLIVINDSPSGDLKQFLTEYSVDNRKIL 63
DB 5 SVMPEVNSE-KYLESESTELNOSYSDLEFIINDSGTNSFK-TKEVAKLDPKRLNV- 61
QY 64 LNEENIGLASSLNKAVKISKEGYIFRMDADISYPSFQKQIRMEEN-SLDSEATLIEL 122

DB 62 ISRENKGIYSLNEAIRLANGEYIARMDADISAPRIEKQISFLKSHNDILGTQVKY 121
QY 123 IDQGNLVYQRESNKNILYNDI-----RKLLNR-SILAPTWCVKKVFDKLMGYRDV 177
DB 122 VGNISNDI-KEKENKLNINFDIYDNRKELIYVWCLNHPSTWFKDILRELKGYNDF- 179
QY 178 PVEDYDPAIGALADPKIGLGNKVLQYRLNENGISQTNKFOQYISALIQDFYKESYI 237
DB 180 KSEDLDMRLAISGCFKLYLKEELIYFRWHEESKTRVDN-----QNEGKDG 229
QY 238 DITKITNYPQEVYIKKRY-----TQOELSKYFE 265
DB 230 KIKLIDVFKRPFKQDFKITVWGASNGKTKYEVLDPEFE 268

RESULT 3

US-10-282-122A-51835

/ Sequence 51835, Application US/10282122A
/ Publication No. US20040029129A1
/ GENERAL INFORMATION:
/ APPLICANT: Wang, Liangsu
/ APPLICANT: Zamudio, Carlos
/ APPLICANT: Malone, Cheryl
/ APPLICANT: Haselbeck, Robert
/ APPLICANT: Ohlsen, Kari
/ APPLICANT: Zyskind, Judith
/ APPLICANT: Wall, Daniel
/ APPLICANT: Trawick, John
/ APPLICANT: Carr, Grant
/ APPLICANT: Yamamoto, Robert
/ APPLICANT: Forsyth, R.
/ APPLICANT: Xu, H.
/ TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
/ FILE REFERENCE: EUTRA.034A
/ CURRENT APPLICATION NUMBER: US/10/282,122A
/ PRIOR FILING DATE: 2003-02-20
/ PRIOR APPLICATION NUMBER: 60/191,078
/ PRIOR FILING DATE: 2000-03-21
/ PRIOR APPLICATION NUMBER: 60/206,848
/ PRIOR FILING DATE: 2000-05-23
/ PRIOR APPLICATION NUMBER: 60/207,727
/ PRIOR FILING DATE: 2000-05-26
/ PRIOR APPLICATION NUMBER: 60/230,335
/ PRIOR FILING DATE: 2000-09-06
/ PRIOR APPLICATION NUMBER: 60/230,347
/ PRIOR FILING DATE: 2000-09-09
/ PRIOR APPLICATION NUMBER: 60/242,578
/ PRIOR FILING DATE: 2000-10-23
/ PRIOR APPLICATION NUMBER: 60/253,625
/ PRIOR FILING DATE: 2000-11-27
/ PRIOR APPLICATION NUMBER: 60/257,931
/ PRIOR FILING DATE: 2000-12-22
/ PRIOR APPLICATION NUMBER: 60/267,636
/ PRIOR FILING DATE: 2001-02-09
/ PRIOR APPLICATION NUMBER: 60/269,308
/ PRIOR FILING DATE: 2001-02-16
/ Remaining Prior Application data removed - See File Wrapper or PALM.
/ NUMBER OF SEQ ID NOS: 78614
/ SOFTWARE: PatentIn version 3.1
/ SEQ ID NO 51835
/ LENGTH: 336
/ TYPE: PRT
/ ORGANISM: Clostridium acetobutylicum
US-10-282-122A-51835

Query Match 21.9%, Score 348.5, DB 12, Length 336;
Best Local Similarity 36.6%, Pred. No. 3, Se-23;
Matches 90, Conservative 46, Mismatches 95, Indels 15, Gaps 8;
QY 4 STMSVNEPLNVRDSESTINOTLDPFEFLIVINDSPSGDLKQFLTEYSVDNRKIL 63
DB 8 SVMPEVNSE-KYLESESTELNOSYSDLEFIINDSGTNSFK-TKEVAKLDPKRLNV- 64

QY 64 LNEENIGLASSLNKAVKISKEGYIFRMDADDISYSPRFDKQIRMEEN-SDPSATLIEL 122
DB 65 ISNEHGLVDLSNEGINARGLTANMDADDISINNRIEKEQFEFLINVDLIGTRIBA 124
QY 123 ---IDQGNLVYKQRESNKIYLTNDIRKMLNRSILAHPTWCYKVKYFDKLMGYR-DLVP 178
DB 125 FGIDEREKQKTIYNSAFSIR-FDSQNIQVFLTSCALPHPSVMEKOSIVKLGKRYREYDT 183
QY 179 VEDYDFAIRGALADFKIGLNLKVLQYRLNENGISQTNKFKQIYSAILODFYKESYID 238
DB 184 AEDYDLMLRALIRNGYKIVRDECLIKRYNNSKTAIVEMPRQVBE-----YTMAXKD 237
QY 239 ITXITN 244
DB 238 YINDTN 243

RESULT 4

US-10-282-122A-52169
Sequence 52169, Application US/10282122A
Publication No. US20040029129A1
GENERAL INFORMATION:
APPLICANT: Wang, Liangsu
APPLICANT: Zamudio, Carlos
APPLICANT: Malone, Cheryl
APPLICANT: Haselbeck, Robert
APPLICANT: Ohlsen, Karl
APPLICANT: Zykkind, Judith
APPLICANT: Wall, Daniel
APPLICANT: Trawick, John
APPLICANT: Carr, Grant
APPLICANT: Yamamoto, Robert
APPLICANT: Forsyth, R.
APPLICANT: Xu, H.
TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
FILE REFERENCE: EPIITRA.034A
CURRENT APPLICATION NUMBER: US/10/282.122A
CURRENT FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21/2006,848
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: 60/230,335
PRIOR FILING DATE: 2000-09-06
PRIOR APPLICATION NUMBER: 60/230,347
PRIOR FILING DATE: 2000-09-09/2006,578
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR FILING DATE: 2000-10-23
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-12-22/2006,636
PRIOR APPLICATION NUMBER: 60/267,636
PRIOR FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/269,308
PRIOR FILING DATE: 2001-02-16
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 78614
SOFTWARE: PatentIn version 3.1
SEQ ID NO 52169
LENGTH: 333
TYPE: PRT
ORGANISM: Clostridium acetobutylicum
US-10-282-122A-52169

Query Match 19.9%; Score 317; DB 12; Length 333;
Best Local Similarity 32.8%; Pred. No. 2,2e-20;
Matches 83; Conservative 53; Mismatches 91; Indels 26; Gaps 10;
QY 4 STMSVYNEPLNYVDVSSESLNQTLTDFEFLIIVDNPSSRDLKQFLTEYSVVDNRKITL 63

DB 3 SVIMEPTNCE-KYIEESIETLKTQYRDFEFLIIVDNGSNDKSI-DIINKYANDNRIVV 60
QY 64 LNEENIGLASSLNKAVKISKEGYIFRMDADDISYSPRFDKQIRMEEN-SDPSATLIEL 122
DB 61 SRDNNMGVYVLSNGIDIRAKGSYARMDADDIALPERFERQIEYLNKQKQVDILACVYA 120
QY 123 IDQGNLVYKQRESNKIYLTND-----IRKMLNRSILAHPTWCYKVKYFDKLMGYR-D 175
DB 121 F---GDVSRQKLEREHWYVNDLNNSESIESLFENCYIAHPSVMEKOSIVKLGKRYNIN 177
QY 176 LVPEYDFAIRGALADFKIGLNLKVLQYRL-NEGISQT---NKFQIYSAILODF 230
DB 178 YKTEDYDLMLRALIRNGYKIMLEKMKIRLHNSKIHRAEGFSSIRDIIGSLR--EY 235
QY 231 YKEX-----SYI 237
DB 236 VKEKXKLDPSYV 248

RESULT 5

US-09-738-626-3905
Sequence 3905, Application US/09738626
Publication No. US20020197605A1
GENERAL INFORMATION:
APPLICANT: NAKAGAWA, SATOSHI
APPLICANT: MIZOGUCHI, HIROSHI
APPLICANT: ANDO, SEIRO
APPLICANT: HAYASHI, MIKIRO
APPLICANT: OCHITAI, KEIKO
APPLICANT: YOKOI, HARUHIKO
APPLICANT: TATEISHI, NAOKO
APPLICANT: SENOH, AKIHIRO
APPLICANT: IKEDA, MASATO
APPLICANT: OZAKI, AKIO
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REFERENCE: 249-125
CURRENT APPLICATION NUMBER: US/09/738.626
CURRENT FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: JP 99/377484
PRIOR FILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: JP 00/159162
PRIOR FILING DATE: 2000-04-07
PRIOR APPLICATION NUMBER: JP 00/280988
PRIOR FILING DATE: 2000-08-03
NUMBER OF SEQ ID NOS: 7059
SOFTWARE: PatentIn ver. 3.0
SEQ ID NO 3905
LENGTH: 274
TYPE: PRT
ORGANISM: Corynebacterium glutamicum
US-09-738-626-3905

Query Match 15.1%; Score 240.5; DB 9; Length 274;
Best Local Similarity 26.2%; Pred. No. 1.4e-13;
Matches 71; Conservative 72; Mismatches 101; Indels 27; Gaps 10;

QY 10 YNEPLNYVDVSSESLNQTLTDFEFLIIVDNPSSRDLKQFLTEYSVVDNRKITLNEENI 69
DB 17 YDE---YCSQSIKSVEQYEWQVLTVDGAPIDVPEWVKEH---ERKIVCEKTIQ 69
QY 70 GLASSLNKAVKISKEGYIFRMDADDISYSPRFDKQIRMEEN-SDPSATLIELIDQGN 128
DB 70 GTFSTLNNIGIKASDGLILARLSDDLAAPSRLSKQBEFLRNHPYIICVATKTHINEHGK 129
QY 129 LVYKQRESNKIYLTNDIRKMLNRSILAHPTWCYKVKYFDKLMGYR-DLVPEDYDFAIR 167
DB 130 IF---GQSADLPSTQDIRQILLVKNPITHSSVVRKQVVEQIGYSLEWTRSDYELFLR 186
QY 188 ----GALADFKIGLNLKVLQYRLNENGIS-OTNKEQIYSAILODFYKESYIDITXI 242
DB 187 LSKTGA-----ICVYDEBSLSYRIHGGQHSRKTSPKXTW-IILKRBWELASFLKRSYV 240
QY 243 TNFQEVYIKRKYTOELSKYFELKSTPBIT 273

Db 241 RQIFLFI---WYGAQVTRYLGLRKAGFMT 267

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RESULT 6
US-09-816-028A-31
: Sequence 31, Application US/09816028A
: Patent No. US20020042369A1
GENERAL INFORMATION:
APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
TITLE OF INVENTION: National Research Council of Canada
TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
FILE OF INVENTION: Gangliosides and Ganglioside Mimics
FILE REFERENCE: 019633-00011US
CURRENT APPLICATION NUMBER: US/09/816,028A
CURRENT FILING DATE: 2001-03-21/81,213
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 31
LENGTH: 297
TYPE: PRT
ORGANISM: Campylobacter jejuni
FEATURE:
OTHER INFORMATION: beta-1,3 galactosyl transferase from C. jejuni O:10
US-09-816-028A-31

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Query Match      14.9%; Score 237; DB 9; Length 297;
Best Local Similarity 27.8%; Pred. No. 3,3e-13;
Matches      85; Conservative 61; Mismatches 112; Indels 48; Gaps 14

Dd      85; Conservative 61; Mismatches 112; Indels 48; Gaps 14

QY      4 SIIMSVYNEPLNYVRDSVESILNQTLTTPFEFIVIDNBSRGDLKQPLTEYSVVDNRKIL 63
      5 SIILPFTYVNE-QYIADIAESCINQTFKNE-IIVDDCGSDKSIDIVKAYAKDDBRKII 62
      6 LNENINIGLASSLNKAKYIKSKGEIIFPMDDDISYRSRDXKDIRMEENSDD--FSATII 120
      63 HNEENILTLARAYEGVKVANSPIYINFLDDDDYLDLEINACEBOMKILKONNEIDLFFNAPVL 122
      121 E--LIDQKNLVYKQRESNKIYLTINDIRKMLNRSIIAHPWC--YKAYFPDLKMGYRD 175
      123 ENNNKIRKRLNP-----QEKCYEKDFIKELIKTKNOLFMTVMKAVIKKELLYKANG--- 173
      176 LVPEVDYDFAIRGALADFKIGLINTKYLQYRLNENGISQT-NKFRQYIYSAIIDDIFYEK 234
      174 LISLE-----NAKINMAEDVLLYPLP--INISNTIFHLKSUNLYQVINFSITK 220
      QY      235 SYDIITKITYPQF-----YVKK-----RYQDLSKFEKLSPIISIRKLYIC-- 280
      Db      221 T-LTLQNIKNTIOBQDNLVILKMKQYNFNMLTLKLIEFLLIEKISLSKRNVLCEK 279
      QY      281 LYLEYK 286
      Db      280 INIEFK 285

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RESULT 7
US-10-303-161-31
; Sequence 31, Application US/10303161
; Publication No. US20030148459A1
; GENERAL INFORMATION:
; APPLICANT: Gilbert, Michel
; APPLICANT: Wakarchuk, Warren W.
; APPLICANT: National Research Council of Canada
; TITLE OF INVENTION: Camphylolactone Glycosyltransferase for Biosynthesis of
; TITLE OF INVENTION: Gangliosides and Ganglioside Mimics
; FILE REFERENCE: 019633-00011IUS
; CURRENT APPLICATION NUMBER: US/10/303,161
; CURRENT FILING DATE: 2002-11-21

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PRIORITY APPLICATION NUMBER: US/09/816,028
PRIORITY FILING DATE: 2001-03-21
PRIORITY APPLICATION NUMBER: US 60/118,213
PRIORITY FILING DATE: 1999-02-01
PRIORITY APPLICATION NUMBER: US 09/495,406
PRIORITY FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 31
LENGTH: 297
TYPE: PRT
ORGANISM: Campylobacter jejuni
FEATURE:
OTHER INFORMATION: beta-1,3 galactosyl transferase from C. jejuni O:10
US-10-303-161-31

```

[illegible]

```

RESULT 8
US-10-303-118-31
/ Sequence 31, Application US/10303118
/ Publication No. US20030157655A1
/
GENERAL INFORMATION:
/
APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
APPLICANT: National Research Council of Canada
/ TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
/ TITLE OF INVENTION: Gangliosides and Ganglioside Mimics
FILE REFERENCE: 019633-000111US
CURRENT APPLICATION NUMBER: US/10/303,118
CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOFTWARE: PatentIn Ver. 2.1
/
/ SEQ ID NO 31
/
/ LENGTH: 297
/
/ TYPE: prt
/
/ ORGANISM: Campylobacter jejuni
/
/ FEATURE:
/
/ OTHER INFORMATION: beta-1,3 galactosyl transferase from C. jejuni O:10
US-10-303-118-31

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RESULT 11
US-10-303-162-31
Sequence 31, Application US/10303162
Publication No. US20030157658A1
GENERAL INFORMATION:
APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
APPLICANT: National Research Council of Canada
TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
TITLE OF INVENTION: Gangliosides and Ganglioside Mimics
FILE REFERENCE: 019633-000111US
CURRENT APPLICATION NUMBER: US/10/303,162
PRIOR FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 31
LENGTH: 297
TYPE: PRF
ORGANISM: Campylobacter jejuni
FEATURE:
OTHER INFORMATION: beta-1,3 galactosyl transferase from C. jejuni O:110
US-10-303-162-31

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Best Local Similarity 27.8%; Pred. No. 3,3e-13;
Matches 85; Conservative 61; Mismatches 112; Indels 48; Gaps 14;
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DB 63 HNEENKILARAREYGKAVNSPYIMFLDPDYLLENACECEKILKNEDILFPNAYVL 122
QY 121 E---LIDQGNLYVKQRESNKIYLINDIRKMLNRSILAHPTWC--VKKKVFDEKMGYRD 175
DB 123 ENNNKIERKLNFE---CEKCYVKCDPKEKELKTKMLTWAKVKKELKXAVG--- 173
QY 176 LVEVEYDPAIRGALADPKIGLNLKYLQYRLNENGISQTNKPKQYISALIDPFYKEX 234
DB 174 LLSLE-----NAKINMAEDVLLYPL--INISYTHLSKMLNYQVNNFSITK 220
QY 235 SYIDIRKINNYQE---YVIRK-----RYTQGLSKYFELKSTPSITRKLKLYIC-- 280
DB 221 T-LTQNIKINIQEQDNVLYLLEKQYNNVNFLLTLKLEYFLLEKXVLSLSKRVLCFK 279
QY 281 LVIYFK 286
DB 280 INIFPK 285

RESULT 12
US-09-816-028a-27
Sequence 27, Application US/09816028A
Patent No. US20020042369A1
GENERAL INFORMATION:
APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
APPLICANT: National Research Council of Canada
TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
TITLE OF INVENTION: Gangliosides and Ganglioside Mimics
FILE REFERENCE: 019633-000111US
CURRENT APPLICATION NUMBER: US/09/816,028A
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01

PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 27
LENGTH: 301
TYPE: PRF
ORGANISM: Campylobacter jejuni
FEATURE:
OTHER INFORMATION: beta-1,3-galactosyltransferase from C. jejuni strain
OTHER INFORMATION: OH4384 (ORF 6a of 11pooligosaacharide (LOS))
OTHER INFORMATION: biosynthesis locus
US-09-816-028a-27

Query Match 14.7%; Score 234; DB 9; Length 301;
Best Local Similarity 28.2%; Pred. No. 6,3e-13;
Matches 88; Conservative 53; Mismatches 135; Indels 36; Gaps 13;
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DB 5 SIILPTVYNE-QYIAAIESCINQTFKIDIE-IIVDDCGNDSINAKESKDKRIKII 62
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DB 63 HNEENKILARAREYGKAVNSPYIMFLDPDYLLENACECEKILKNEDILFPNAYVL 121
QY 124 DQGNLYVKQRESNKIYLINDIRKMLNRSILAHPTWC--VKKKVFDEKMGYRDLPVE 180
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QY 181 DYDFAIRGALADPKIGLNLKYLQYRLNENGISQTNKPKQYISALIDPFYKEXYIDIT 240
DB 172 ---FASLRREKQDKINMAEDVLLYPM---LSQAQKIA--YVNCNLVYVPPNNNSICNT 222
QY 241 K---ITNYQEVVKKRYQGE--LSKYFELKSTPSITIR-KLYICLVLYKSPVLRSL 293
DB 223 KNEVLYKNQIQEQLVNINRYQVYILNKC--SVLYLILKYLVIQYIKIRKTLMTVL 279
QY 294 LINDINILVLEKL 305
DB 260 LAK-INILTLKI 290

RESULT 13
US-10-303-161-27
Sequence 27, Application US/10303161
Publication No. US20030148459A1
GENERAL INFORMATION:
APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
APPLICANT: National Research Council of Canada
TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
TITLE OF INVENTION: Gangliosides and Ganglioside Mimics
FILE REFERENCE: 019633-000111US
CURRENT APPLICATION NUMBER: US/10/303,161
PRIOR FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 27
LENGTH: 301
TYPE: PRF
ORGANISM: Campylobacter jejuni
FEATURE:
OTHER INFORMATION: beta-1,3-galactosyltransferase from C. jejuni strain
OTHER INFORMATION: OH4384 (ORF 6a of 11pooligosaacharide (LOS))
OTHER INFORMATION: biosynthesis locus
US-10-303-161-27

Query Match 14.7%; Score 234; DB 14; Length 301;
 Best Local Similarity 28.2%; Pred. No. 6.3e-13;
 Matches 88; Conservative 53; Mismatches 135; Indels 36; Gaps 13;

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US-10-303-118-27
 ; Sequence 27, Application US/10303118
 ; Publication No. US20030157656A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Wakarchuk, Michel
 ; APPLICANT: National Research Council of Canada
 ; TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
 ; FILE OF INVENTION: Gangliosides and Ganglioside Mimics
 ; FILE REFERENCE: 019633-000111US
 ; CURRENT APPLICATION NUMBER: US/10/303,118
 ; CURRENT FILING DATE: 2002-11-21
 ; PRIOR APPLICATION NUMBER: US/09/816,028
 ; PRIOR FILING DATE: 2001-03-21
 ; PRIOR APPLICATION NUMBER: US 60/118,213
 ; PRIOR FILING DATE: 1999-02-01
 ; PRIOR APPLICATION NUMBER: US 09/495,406
 ; PRIOR FILING DATE: 2000-01-31
 ; NUMBER OF SEQ ID NOS: 49
 ; SOFTWARE: PatentIn Ver. 2.1
 ; SEQ ID NO 27
 ; LENGTH: 301
 ; TYPE: PRF
 ; ORGANISM: Campylobacter jejuni
 ; FEATURES:
 ; OTHER INFORMATION: beta-1,3-galactosyltransferase from C. jejuni strain
 ; OTHER INFORMATION: OH4384 (ORF 6a of lipooligosaccharide (LOS))
 ; OTHER INFORMATION: biosynthesis locus)
 US-10-303-118-27

Query Match 14.7%; Score 234; DB 14; Length 301;
 Best Local Similarity 28.2%; Pred. No. 6.3e-13;
 Matches 88; Conservative 53; Mismatches 135; Indels 36; Gaps 13;
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 ; Sequence 27, Application US/10303128
 ; Publication No. US20030157656A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Wakarchuk, Michel
 ; APPLICANT: National Research Council of Canada
 ; TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
 ; FILE OF INVENTION: Gangliosides and Ganglioside Mimics
 ; FILE REFERENCE: 019633-000111US
 ; CURRENT APPLICATION NUMBER: US/10/303,128
 ; CURRENT FILING DATE: 2002-11-21
 ; PRIOR APPLICATION NUMBER: US/09/816,028
 ; PRIOR FILING DATE: 2001-03-21
 ; PRIOR APPLICATION NUMBER: US 60/118,213
 ; PRIOR FILING DATE: 1999-02-01
 ; PRIOR APPLICATION NUMBER: US 09/495,406
 ; PRIOR FILING DATE: 2000-01-31
 ; NUMBER OF SEQ ID NOS: 49
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 ; SEQ ID NO 27
 ; LENGTH: 301
 ; TYPE: PRF
 ; ORGANISM: Campylobacter jejuni
 ; FEATURES:
 ; OTHER INFORMATION: beta-1,3-galactosyltransferase from C. jejuni strain
 ; OTHER INFORMATION: OH4384 (ORF 6a of lipooligosaccharide (LOS))
 ; OTHER INFORMATION: biosynthesis locus)
 US-10-303-128-27

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Page 8

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Mon Mar 29 10:18:58 2004

us-09-900-038a-2.rmpb

Page 1

GenCore version 5.1.6
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Published Applications NA:*

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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

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3	118	12.6	1002	12 US-10-282-122a-15868	Sequence 15868, A
4	95.2	10.1	1011	12 US-10-282-122a-15851	Sequence 15851, A
5	84	8.9	1002	12 US-10-282-122a-15855	Sequence 15855, A
6	76.2	8.1	3673778	14 US-10-312-841-1	Sequence 1, Appl
7	73.6	7.8	1830121	14 US-10-329-960-1	Sequence 1, Appl
8	73.6	7.8	1830121	15 US-10-329-960-1	Sequence 1, Appl
9	73.2	7.8	891	9 US-09-816-028a-30	Sequence 30, Appl
10	73.2	7.8	891	14 US-10-303-118-30	Sequence 30, Appl
11	73.2	7.8	891	14 US-10-303-118-30	Sequence 30, Appl
12	73.2	7.8	891	14 US-10-303-128-30	Sequence 30, Appl
13	73.2	7.8	891	14 US-10-303-134-30	Sequence 30, Appl
14	73.2	7.8	891	14 US-10-303-162-30	Sequence 30, Appl
15	71.6	7.6	747	12 US-10-282-122a-17153	Sequence 17153, A

C 16	70.2	7.5	15548	14 US-10-311-455-2128	Sequence 2128, Ap
C 17	69.8	7.4	14187	14 US-10-114-170-121	Sequence 121, Ap
C 18	69.4	7.4	158001	16 US-10-211-179-11	GENERAL INFORMAT
C 19	68.6	7.3	5371	14 US-10-311-455-2267	Sequence 2267, Ap
C 20	67	7.1	4915	9 US-09-070-927a-125	Sequence 125, Ap
C 21	64.8	6.9	906	9 US-09-816-028a-26	Sequence 26, Appl
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C 32	64.8	6.9	11474	14 US-10-303-162-1	Sequence 1, Appl
C 33	64	6.8	7597	14 US-10-311-455-986	Sequence 986, App
C 34	63.8	6.8	1170	9 US-09-816-028a-15	Sequence 15, Appl
C 35	63.8	6.8	1170	14 US-10-303-161-15	Sequence 15, Appl
C 36	63.8	6.8	1170	14 US-10-303-118-15	Sequence 15, Appl
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C 42	63.8	6.8	11474	14 US-10-303-118-1	Sequence 1, Appl
C 43	63.8	6.8	11474	14 US-10-303-128-1	Sequence 1, Appl
C 44	63.8	6.8	11474	14 US-10-303-134-1	Sequence 1, Appl
C 45	63.8	6.8	11474	14 US-10-303-162-1	Sequence 1, Appl

ALIGNMENTS

RESULT 1
US-09-900-038a-2
; Sequence 2, Application US/09900038A
; Patent No. US20020142425A1

GENERAL INFORMATION:
; APPLICANT: Miyake, Katsunide
; APPLICANT: Matsubae, Masaki
; APPLICANT: Iijima, Shintu
; TITLE OF INVENTION: Beta 1,3-galactosyltransferase and DNA encoding the same
; FILE REFERENCE: 766.53
; CURRENT APPLICATION NUMBER: US/09/900,038A
; CURRENT FILING DATE: 2001-09-21
; PRIOR APPLICATION NUMBER: JP 2001-392
; PRIOR FILING DATE: 2001-01-05
; NUMBER OF SEQ ID NOS: 8
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 2
; TYPE: DNA
; ORGANISM: Streptococcus agalactiae Type Ib
US-09-900-038a-2

Query Match 100.0%; Score 939; DB 9; Length 939;
Best Local Similarity 100.0%; Pred. No. 3.2e+166;
Matches 939; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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DB 121 CCAAGTAGAGGATTTAAGCAATTTCTTAACAGATATTCAGTTGTGATATGAGATTA 180

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DB 901 TTAGTACTGAAATTTGTTGAGAGAGAGAAACAAAGTGAC 939

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RESULT 2
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; Sequence 3, Application US/09900038a
; Patent No. US2002042425A1
; GENERAL INFORMATION:
; APPLICANT: Miyake, Katsuhide
; APPLICANT: Matanabe, Masaki
; APPLICANT: Iijima, Shinji
; TITLE OF INVENTION: Beta 1,3-galactosyltransferase and DNA encoding the same
; FILE REFERENCE: 766.53
; CURRENT APPLICATION NUMBER: US/09/900,038a
; PRIOR FILING DATE: 2001-09-21
; PRIOR APPLICATION NUMBER: JP 2001-392
; NUMBER OF SEQ ID NOS: 8
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 3
; LENGTH: 6865
; TYPE: DNA
; ORGANISM: Streptococcus agalactiae type Ib

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FEATURE:
NAME/KEY: CDS
LOCATION: (617)..(1789)
NAME/KEY: CDS
LOCATION: (1816)..(2262)
NAME/KEY: CDS
LOCATION: (2265)..(2744)
NAME/KEY: CDS
LOCATION: (2943)..(3979)
NAME/KEY: CDS
LOCATION: (3982)..(4953)
NAME/KEY: CDS
LOCATION: (5009)..(5947)
US-09-900-038a-3

Query Match      100.0%; Score 939; DB 9; Length 6865;
Best Local Similarity 100.0%; Pred. No. 6, 4e-166;
Matches 939; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ATGAATTAATAGTATCATTAATGTCGTATATATATGAGCTTTAAATTAATGAGAGATTCA 60
DB 5009 ATGAATTAATAGTATCATTAATGTCGTATATATATGAGCTTTAAATTAATGAGAGATTCA 5068
QY 61 GTAAATCTAATTAATTAATCAAGCTTACTGATTTGAGTCAATTAATGTCATTGATTAAT 120
DB 5069 GTAAATCTAATTAATTAATCAAGCTTACTGATTTGAGTCAATTAATGTCATTGATTAAT 5128
QY 121 CCAAGTAGAGGTGATTTTAAAGCAATCTTAACAGAAATTCAGTTGATTAATTAAGATA 180
DB 5129 CCAAGTAGAGGTGATTTTAAAGCAATCTTAACAGAAATTCAGTTGATTAATTAAGATA 5188
QY 181 AAAATCTGCTTAATGAGAGAAATATGCTTTGAGCATCAAGTTGAACAAAGGGTGA 240
DB 5189 AAAATCTGCTTAATGAGAGAAATATGCTTTGAGCATCAAGTTGAACAAAGGGTGA 5248
QY 241 ATTTCTAAGGAGATATATTTTGAATGAGATGCTGATGATATTTCAATCCAGTGA 300
DB 5249 ATTTCTAAGGAGATATATTTTGAATGAGATGCTGATGATATTTCAATCCAGTGA 5308
QY 301 TTTGATAGCAAAATTCGTTTATGAGAGAAATTCATGATTTTTCAGCACTTAATA 360
DB 5309 TTTGATAGCAAAATTCGTTTATGAGAGAAATTCATGATTTTTCAGCACTTAATA 5368
QY 361 GAATTTAGAGCCAAAGAGAAATTTAGTATTAACAAGAGAAATTAATAATTAATAATTA 420
DB 5369 GAATTTAGAGCCAAAGAGAAATTTAGTATTAACAAGAGAAATTAATAATTAATAATTA 5428
QY 421 TTAATTAATGATATGAGAGAGATGTTTGAATGATCTTACTGCCCCACCAAGCTGG 480
DB 5429 TTAATTAATGATATGAGAGAGATGTTTGAATGATCTTACTGCCCCACCAAGCTGG 5488
QY 481 TGGGTAAAAAGAAAGTTTTCGTTTATGAGAGAAATTCATGAGATTTCTCAGCACTTA 540
DB 5489 TGGGTAAAAAGAAAGTTTTCGTTTATGAGAGAAATTCATGAGATTTCTCAGCACTTA 5548
QY 541 GATTATGATTTTGAATTAAGAGAGCTCTGGCTGATTTCAAAATCGCTTACTCAATTA 600
DB 5549 GATTATGATTTTGAATTAAGAGAGCTCTGGCTGATTTCAAAATCGCTTACTCAATTA 5608
QY 601 GTACTTTTACAGTATGATTTTGAAGAGATGGAATTCACAAACCAATTAAGTTAAAGCA 660
DB 5609 GTACTTTTACAGTATGATTTTGAAGAGATGGAATTCACAAACCAATTAAGTTAAAGCA 5668
QY 661 TATATTTACTCAGCTATTTTGAAGAGATTTTGAAGAGATTTTGAAGAGATTTTGAAGATTA 720
DB 5669 TATATTTACTCAGCTATTTTGAAGAGATTTTGAAGAGATTTTGAAGAGATTTTGAAGATTA 5728
QY 721 AAAATTTACTAATTTCTTCAAGAGATGTAATAAGAAAGCTTACTCAGCAAGAGCTC 780
DB 5729 AAAATTTACTAATTTCTTCAAGAGATGTAATAAGAAAGCTTACTCAGCAAGAGCTC 5788
QY 781 TCTAAATATTTGAGCTAAATTTGAGAGAGCTCTGGCTGATTTCAAAATCGCTTACTCAATTA 840
DB 781 TCTAAATATTTGAGCTAAATTTGAGAGAGCTCTGGCTGATTTCAAAATCGCTTACTCAATTA 840

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Db	5789	TCCTAATATTCTTGAGCTAAATCTCCCTCGATATGATATGAGAAACINAAITTTGT	5848
Qy	841	TTAATATTATCTTTAAAGTCTCCCTGGTGTAGAGAGTATTAATAAATGATTTATATT	900
Db	5849	TTATATTATACCTTTAACTCTCCCTGGTGTAGAGAGTATTATATAAGATTTATATT	5900
Qy	901	TTAGTACTGAATTTGTTTGGAGAGAGAAACAAAGTGAC	939
Db	5909	TTAGTACTGAATTTGTTTGGAGAGAGAAACAAAGTGAC	5947

RESULT 3
ITS-10-28

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1  / Sequence 1988, Application US/10289122A
2  / Publication No. US20040029129A1
3  / GENERAL INFORMATION:
4  / APPLICANT: Wang, Liangsu
5  / APPLICANT: Zamudio, Carlos
6  / APPLICANT: Malone, Cheryl
7  / APPLICANT: Haselbeck, Robert
8  / APPLICANT: Ohlsen, Kari
9  / APPLICANT: Zykkind, Judith
10 / APPLICANT: Wall, Daniel
11 / APPLICANT: Trawick, John
12 / APPLICANT: Carr, Grant
13 / APPLICANT: Yamamoto, Robert
14 / APPLICANT: Forsyth, R.
15 / APPLICANT: Xu, H.
16 / TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
17 / FILE REFERENCE: ELITRA.024A
18 / CURRENT APPLICATION NUMBER: US/10/0282.122A
19 / CURRENT FILING DATE: 2003-02-20
20 / PRIOR APPLICATION NUMBER: 60/191,078
21 / PRIOR FILING DATE: 2000-03-21
22 / PRIOR APPLICATION NUMBER: 60/206,848
23 / PRIOR FILING DATE: 2000-05-23
24 / PRIOR APPLICATION NUMBER: 60/207,727
25 / PRIOR FILING DATE: 2000-05-26
26 / PRIOR APPLICATION NUMBER: 60/230,335
27 / PRIOR FILING DATE: 2000-09-06
28 / PRIOR APPLICATION NUMBER: 60/230,347
29 / PRIOR FILING DATE: 2000-09-09
30 / PRIOR APPLICATION NUMBER: 60/242,578
31 / PRIOR FILING DATE: 2000-10-23
32 / PRIOR APPLICATION NUMBER: 60/253,625
33 / PRIOR FILING DATE: 2000-11-27
34 / PRIOR APPLICATION NUMBER: 60/257,931
35 / PRIOR FILING DATE: 2000-12-22
36 / PRIOR APPLICATION NUMBER: 60/267,636
37 / PRIOR FILING DATE: 2001-02-09
38 / PRIOR APPLICATION NUMBER: 60/269,308
39 / PRIOR FILING DATE: 2001-02-16
40 / Remaining Prior Application data removed - See File Wrapper or PALM.
41 / NUMBER OF SEQ ID NOS: 78614
42 / SOFTWARE: PatentIn version 3.1
43 / SEQ ID NO 15886
44 / LENGTH: 1002
45 / TYPE: DNA
46 / ORGANISM: Clostridium acetobutylicum
47 / US-10-282-122A-15886

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Query Match 12.6%; Score 118; DB 12; Length 1002;
Best Local Similarity 50.1%; Pred. No. 1,2e-12;
Matches 467; Conservative 0; Mismatches 440; Indels 25; Gaps 6

QY 13 AHCATATGTCGGTATATATGACCTTTAAATAGTAGAGATTCAGAGAACTATTA 72
Db 13 AGCGTTGTATGCGCACTTTATATATAGTAAAAATCTGAAGAGTCCTAGAAATATT 72
QY 73 TTAAATCAACGCTTACTGATTTTGAGTTCTATATGTCTATTATATCCAAGTAGAG 132
Db 73 TTAAATCAATCATACAGTGAATTTTGGAATTTATATATATATATATAGTGCCTC--TACGAT 129

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QY	133	GAATTAAGACATTCCTTACAGAAATATTCAGTTGTAATATAGATTAATAAATCTTGCTT	192
Db	130	AACCTCTTTAAATATATTAAAGAAATATCCAAATTAGATTAAGAATATAA---TGTGATT	186
QY	193	AATGAAAGAAATATTTGGTTTAGCATCAAGTTTGAACCAAGCGGTGAAAATTTCTAAAGCA	252
Db	187	TCAAAGGAAAAATAAGGAATTGTGTACTCGTTAATATGAACTATACGATTGGCAAAAGCA	246
QY	253	GAATATATTTTGAATGCGATGATGATATTTTCATATCCAGTGAATTTGATAAGCA	312
Db	247	GAATATATATGCAAGATGATGCTGATGATATTTTCGCCACCCAAAGAAATGAAAAACAA	306
QY	313	ATTCTGTTTATGGA---GGAATATTCATGGATTTCTCAGCAACTGTAAATGAAATGATA	369
Db	307	ATTCTCTTTCTAAAAAGCCACAGATATATGATATCTGGGAACACAGTAAGAAGTGATG	366
QY	370	GACCAAAAAAGAAATTTAGTATATTAACACAGAAAGTAATATAA-----TA	417
Db	367	GGCAATATTTCTAAAGATATATAAGAAAAAATATGAATATAGCTAAATATATGATTTGAT	426
QY	418	TACTTACATATGATATATGCAAGAGTGTATATGATATGATCTATCTGCGCCACCCAAAG	477
Db	427	ATTATATGATATATATGAGAAAAATATTTAATTATTTGATTTAGTACATCTTTTA	486
QY	478	TGGTGGCTAAAAAAGAAAGTTTTCGATATAGTTAATGGAATATAGAGATTTAGTACTGTT	537
Db	487	GTATGTGTTTAGGAAGATATATTTAGAGAGCTAAAGATACATATGATTTTAATC---A	543
QY	538	GAAGATTAATGATTTTGCAATTAAGAGAGCTCTGGCTGATTTCAAAATCGGCTTATCTAAT	597
Db	544	GAGATTTTATGATTTATGCTATAGAGCTATTAAGAAAGCGAATTTTAAAAATATTAAGCTTAAG	603
QY	598	AAGATCTTTACAGATATGATTAACAGAAATGGAATATACAAACCATTAAGTTTAAG	657
Db	604	GAAAGACTTATATATTTTATTTGATGCTGCAGAGAGTCAAAGACTAGATATGATTAATCAAAT	663
QY	658	CAATATATTTACTACGCTATTTTAAATATTTTATTAAGAAAAATCTTATATGATATATC	717
Db	664	TATGAAAGTTTAAAGATGGAATATAAATAATTAATATAGATGTTTTAAGAGAGAGTTTC	723
QY	718	ACAAAATATCTATATCTTCTAAGAGATATGATTAAGAAAGCCCTATCTACAGCAAGG	777
Db	724	AAGAAAGACTTAAATAATCACTAGTATAGGGGTCCAGATTAAGAGGCCAAAAATTAACAAAGAG	783
QY	778	CTCTCTAAATATTTAGCTAAATATCTACCCCTGATATTAATATTAAGAAACTATATAT	837
Db	784	GTTTATAGCAATTTTATGAGAAATCTCAATGATATGATTTGTATGATTAATTTTAAACA	843
QY	838	TGTTATATTTATATCTTAAAGTCTCC-CTTGGTTTAGAGGTTATTTAATTAATGATATTA	886
Db	844	GGTAAATTTGAAAAAATTTAGAAATTTCTATCTCAAAAGATATTAATCAATATTAATTTGAT	903
QY	897	TATTTATAGTACGAATTTGTTGGAGGAGAG	928
Db	904	TATGTTTATATGCAACTAACCCGGGAAAAG	935

RESULT 4

Sequence 15651, Application US/10289122A1
Publication No. US20040029129A1
GENERAL INFORMATION:
APPLICANT: Wang, Liangsu
APPLICANT: Zamudio, Carlos
APPLICANT: Malone, Cheryl
APPLICANT: Haselbeck, Robert
APPLICANT: Ohlsen, Kari
APPLICANT: Zyskind, Judith
APPLICANT: Wall, Daniel
APPLICANT: Trewick, John
APPLICANT: Carr, Grant
APPLICANT: Yamamoto, Robert
APPLICANT: Foreyth, R.

APPLICANT: Xu, H.
 TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
 FILE REFERENCE: ELITRA.034A
 CURRENT APPLICATION NUMBER: US/10/282,122A
 CURRENT FILING DATE: 2003-02-20
 PRIOR APPLICATION NUMBER: 60/191,078
 PRIOR FILING DATE: 2000-03-21
 PRIOR APPLICATION NUMBER: 60/206,848
 PRIOR FILING DATE: 2000-05-23
 PRIOR APPLICATION NUMBER: 60/207,727
 PRIOR FILING DATE: 2000-05-26
 PRIOR APPLICATION NUMBER: 60/230,335
 PRIOR FILING DATE: 2000-09-06
 PRIOR APPLICATION NUMBER: 60/230,347
 PRIOR FILING DATE: 2000-09-09
 PRIOR APPLICATION NUMBER: 60/242,578
 PRIOR FILING DATE: 2000-10-23
 PRIOR APPLICATION NUMBER: 60/253,625
 PRIOR FILING DATE: 2000-11-27
 PRIOR APPLICATION NUMBER: 60/257,931
 PRIOR FILING DATE: 2000-12-22
 PRIOR APPLICATION NUMBER: 60/267,636
 PRIOR FILING DATE: 2001-02-09
 PRIOR APPLICATION NUMBER: 60/269,308
 PRIOR FILING DATE: 2001-02-16
 Remaining Prior Application data removed - See File Wrapper or PALM.
 NUMBER OF SEQ ID NOS: 78614
 SOFTWARE: PatentIn version 3.1
 SEQ ID NO: 15651
 LENGTH: 1011
 TYPE: DNA
 ORGANISM: Clostridium acetobutylicum
 US-10-282-122A-15651

Query Match 10.1%; Score 95.2; DB 12; Length 1011;
 Best Local Similarity 50.1%; Pred. No. 2.3e-08;
 Matches 353; Conservative 0; Mismatches 333; Indels 18; Gaps 4;
 Db 46 TATGTAGAGATTCACTAGATATATTAATCAAAAGCTTACGATTTTGAAGTTCTA 105
 55 TATTTGCGAAGCTATAGAAAGTATTTAGATCAGACTATATAGATTTTGAAGTTTATA 114
 106 ATGTGATGATATATCAAGTAGAGGTGATTTAAAGCAATTTCTTAACAGAAATTCAGTT 165
 115 ATAGTGTGATGATGTTCTACTGATGATCTATTA-----CATATATCTAGTACGCT 168
 166 GTGATATATGATATATAAATCTGCTTAATGAAGAAATATGTTTACATCAAGTTTG 225
 169 AATTAAGACAAAGAAATATGATTTCAAGAAACATAGGGGACTTGTGATTCATTA 228
 226 AACAAAGCGGTGAATAATTTCTAAGGAGATATATATTTAGATGAGTCTGATGATAT 285
 229 AATGAAGGTATTAACATAGCTAGAGAAATATACATGACAAAGATGATCGGATGATATA 288
 286 TCAATTCAGATGATTTGATTAAGCAAAATTCGTTTATGAGAGAAATTC---TTGAT 342
 289 TCAATTAATATAGAAATGAAAAACAAATTTGAGTTTATGAATTAACAAAGATGATGAT 348
 343 TTCTCAGCACTCTAATAGATTTGATAGCAAAAAG-----GAATTTAGATATATA 396
 349 ATATTAAGTACTAGAAATAGAGGCTTTGAGATATAGATGAGAAGCAAAAACAAATTAAT 408
 397 CAACGAGAAAGTATATAATATCTTACTATAGATAGATAGAGAAATGTTTATGATATA 456
 409 AATAGGCACTTTTCATTAATAATTTGATTCACAAAATATAGACAGGTTTATTAACCTCA 468
 457 TCTATCTGCCCCCAACGCTGCGTAAATAAAGAAAGTTTTCATTAAGTTATGGA 516
 469 TGTGCTATACCTACTCTTCACTGATGTTTAAAGAGATAGATATCGTAATATTAAGGGA 528
 517 TATAG---AGATTTAGTACTGTTGAAGATTTATGATTTTGCATTAAGAGAGCTCGGCT 573
 529 TATGAAAGAAATATGATCTGCAAGAGATTAATGATTTGTGTTAAGGCTATTAAGAAAT 588

Qy 574 GATTCAAAATGCGCTTACTCAATAAAGTACTTTACGATATAGATTAACAGAAATGCA 633
 Db 589 GATATATAGATGATGAGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 648
 Qy 634 ATATCAAAACCAATAGTTTAAAGCAATATATTTACTGCTATTTTACAGATTTTAT 693
 Db 649 AAAACAGCGGTGAAATGTTTAACTCTAAGATGTTGAATATACATGAAAGCAAGATA 708
 Qy 694 AAAGAAATCTTATATGATATCAACAAATTTACTTAATCTT 737
 Db 709 GATTATATTAATGATATCAATATAAAGCAAGATGATTAATT 752

RESULT 5
 US-10-282-122A-15985
 Sequence 15985, Application US/10282122A
 Publication No. US20040029129A1
 GENERAL INFORMATION:
 APPLICANT: Wang, Liangsu
 APPLICANT: Zamudio, Carlos
 APPLICANT: Malone, Cheryl
 APPLICANT: Haselbeck, Robert
 APPLICANT: Ohlsen, Karl
 APPLICANT: Zyskind, Judith
 APPLICANT: Wall, Daniel
 APPLICANT: Trawick, John
 APPLICANT: Carr, Grant
 APPLICANT: Yamamoto, Robert
 APPLICANT: Forsyth, R.
 APPLICANT: Xu, H.
 TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
 FILE REFERENCE: ELITRA.034A
 CURRENT APPLICATION NUMBER: US/10/282,122A
 CURRENT FILING DATE: 2003-02-20
 PRIOR APPLICATION NUMBER: 60/191,078
 PRIOR FILING DATE: 2000-03-21
 PRIOR APPLICATION NUMBER: 60/206,848
 PRIOR FILING DATE: 2000-05-23
 PRIOR APPLICATION NUMBER: 60/207,727
 PRIOR FILING DATE: 2000-05-26
 PRIOR APPLICATION NUMBER: 60/230,335
 PRIOR FILING DATE: 2000-09-06
 PRIOR APPLICATION NUMBER: 60/230,347
 PRIOR FILING DATE: 2000-09-09
 PRIOR APPLICATION NUMBER: 60/242,578
 PRIOR FILING DATE: 2000-10-23
 PRIOR APPLICATION NUMBER: 60/253,625
 PRIOR FILING DATE: 2000-11-27
 PRIOR APPLICATION NUMBER: 60/257,931
 PRIOR FILING DATE: 2000-12-22
 PRIOR APPLICATION NUMBER: 60/267,636
 PRIOR FILING DATE: 2001-02-09
 PRIOR APPLICATION NUMBER: 60/269,308
 Remaining Prior Application data removed - See File Wrapper or PALM.
 NUMBER OF SEQ ID NOS: 78614
 SOFTWARE: PatentIn version 3.1
 SEQ ID NO: 15985
 LENGTH: 1002
 TYPE: DNA
 ORGANISM: Clostridium acetobutylicum
 US-10-282-122A-15985

Query Match 8.9%; Score 84; DB 12; Length 1002;
 Best Local Similarity 57.9%; Pred. No. 2.8e-06;
 Matches 169; Conservative 0; Mismatches 120; Indels 3; Gaps 1;
 Qy 42 AAATATGTGAGAGATTCACTAGATATATTAATCAAAAGCTTACTGATTTTGAATT 101
 Db 36 AAAATATTTAGAGATATATAGAAAGATATCAATAAAGATATAGATTTTGAATT 95
 Qy 102 CATTAATGTCTATGATATATCAAGTAGAGGTGATTTAAAGCAATTTCTTAACGAATATTC 161

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Db      96 TATTATTGTAATATATGGTTCAAAT---GATAAATCAATGCATTAATTAATTAATTAATGTC 152
Oy      162 AGTGTAGATATATAGAAATTAATAAAATCTTGCTTAATGAGAAAAATATGTTTACATCAAG 221
Db      153 AAATGACGATATATGAATATAGTTGTAGTATCAAGGAGCAACAACATGGGAATGGTGTATTC 212
Oy      222 TTGAAACAAACCGGTGAAAAATTTCTAAGGGAGAAATATTTTTCATGATGATGCTGATGA 281
Db      213 TCTAAATGAGGAAATCGATATAGAGCAAAAAGGAAGTTATATGTCAGCAAAATGATGGCGACGA 272
Oy      282 TATTTCATATCCAAAGTAGATTGATTAAGCAAAATTCGTTTATGAGGAAAAAT 333
Db      273 TATAGCTCTTCTCGAAAGGTTTGAAAGCAATATGATATCTTAATTAATAAT 324

RESULT 6
US-10-312-841-1/c
Sequence 1, Application US/10312841
Publication No. US20030186277A1
GENERAL INFORMATION:
APPLICANT: Epigenome AG
TITLE OF INVENTION: Diagnose von bedeutenden genetischen Parametern innerhalb des KMC
FILE REFERENCE: E01/1208/WO
CURRENT APPLICATION NUMBER: US/10/312,841
CURRENT FILING DATE: 2002-12-30
NUMBER OF SEQ ID NOS: 2
SEQ ID NO 1
LENGTH: 3673778
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
FEATURE:
NAME/KEY: unsure
LOCATION: (3294164)
US-10-312-841-1

Query Match
Best Local Similarity 44.6%; Pred. No. 0.0014;
Matches 341; Conservative 0; Mismatches 422; Indels 1; Gaps 1;

Oy      74 TAAATCAAAACGCTTACTGATTTTGAGTTCATATATGTCATGATATAATCCAGTAGAGGTG 133
Db      1715364 TAAATATATCATATATAAAATATATATATATATTAATTAATATATATATATATATAT 1715305
Oy      134 ATTTAAGCAATTCCTTAACAGATATTCAGTTGTGATATATAGAAATTAATAAAATCTTGCTTA 193
Db      1715304 AATATATTAATATATATATATATATATATATATATATATATATATATATATATATAT 1715245
Oy      194 ATGAGAAAAATTTGGTTTGAATCATCAAGTTGAACAAACGGGTGAAAAATTTCTAAGGAG 253
Db      1715244 AAAATATATATATTAATAATATATATATATATATATATATATATATATATATATAT 1715185
Oy      254 AATATATTTTGAATGATGATGATGATGATATTTTCATATCCAGTAGAGATTGCTAAGCAAA 313
Db      1715184 TATATATATTAATATATATATATATATATATATATATATATATATATATATATATATAT 1715125
Oy      314 TTGCTTTATGAGAAAAATTCATGGAATTTCTCAGCAACCTTAATAGATGATAGACC 373
Db      1715124 TAAATATATATATATATATATATATATATATATATATATATATATATATATATATAT 1715066
Oy      374 AAAAAGGAATTTAGATATATTAACAACGAGAAAGTATATATATATATATATATATATATAT 433
Db      1715065 AAAATATATATATATATATATATATATATATATATATATATATATATATATATATATAT 1715006
Oy      434 TACGGAAGATGTTATTTGAATAGATCTATACTTGCCACCAACGTTGGCGTAAAAAGAA 493
Db      1715005 AATATATATATATATATATATATATATATATATATATATATATATATATATATATATAT 1714946
Oy      494 AAGTTTCGATTAAGTAAATGCGGATATAGAGATTTAAGTACTGTGTAGAGATTTGATTTTG 553
Db      1714945 AAAATATATATATATATATATATATATATATATATATATATATATATATATATATATAT 1714886

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OY      554 CATATAGAGGAGCTCGGTGGCATTTTCAAATPGCGCTTACCAATAAAGTACTTTACGT 613
Db      1714885 TATTATATTAATATATATATATATATATATATATATATATATATATATATATATAT 1714826
OY      614 ATAGATTAAACGAGAAATGAATATCACAAACCAATTAAGTTTAGCAATATATTTACTCAG 673
Db      1714825 ATAAATATATATATATATATATATATATATATATATATATATATATATATATATATAT 1714766
OY      674 CTATTTTACAAGATTTTTTATTAAGAAAAATCTTATATGTGAATACAAAATTTCTAT 733
Db      1714765 TACATATATTAATATATATATATATATATATATATATATATATATATATATATAT 1714706
OY      734 ACTTCAAGATATGTATGATTAAGAAGCGTATTTACTCAGCAAGAGCTCTCTAAATATTTTG 793
Db      1714705 AAATATATCATATTAATAATATATACATATATAAAATATATATATATATATATATATATAT 1714646
OY      794 AGCTTAATATCTAACCCCTAGTATTTACTATTTAGAAAGCTATATTT 838
Db      1714645 ATAAATATATATATATATATATATATATATATATATATATATATATATATATATAT 1714601

RESULT 7
US-10-329-960-1
Sequence 1, Application US/10329960
Publication No. US2003009277A1
GENERAL INFORMATION:
APPLICANT: Fleischmann et al.
TITLE OF INVENTION: Nucleotide Sequence of the Haemophilus influenzae Rd Genome, Fra
FILE REFERENCE: PB186P1
CURRENT APPLICATION NUMBER: US/10/329,960
PRIOR FILING DATE: 2003-01-02
PRIOR APPLICATION NUMBER: US 09/643,990
PRIOR FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: US 08/487,429
PRIOR FILING DATE: 1995-06-07
PRIOR APPLICATION NUMBER: US 08/426,787
PRIOR FILING DATE: 1995-04-21
NUMBER OF SEQ ID NOS: 1
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
LENGTH: 1830121
TYPE: DNA
ORGANISM: Haemophilus influenzae
FEATURE:
NAME/KEY: misc.feature
LOCATION: (4747)..(4747)
OTHER INFORMATION: n equals a, t, g or c
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NAME/KEY: misc.feature
LOCATION: (9921)..(9921)
OTHER INFORMATION: n equals a, t, g or c
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NAME/KEY: misc.feature
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OTHER INFORMATION: n equals a, t, g or c

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NAME/KEY: misc_feature
LOCATION: (40808)..(40810)
OTHER INFORMATION: n equals a, t, g or c
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LOCATION: (44416)..(44416)
OTHER INFORMATION: n equals a, t, g or c
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LOCATION: (44905)..(44905)
OTHER INFORMATION: n equals a, t, g or c
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LOCATION: (44975)..(44975)
OTHER INFORMATION: n equals a, t, g or c
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LOCATION: (45593)..(45593)
OTHER INFORMATION: n equals a, t, g or c
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NAME/KEY: misc_feature
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NAME/KEY: misc_feature
LOCATION: (47036)..(47036)
OTHER INFORMATION: n equals a, t, g or c
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NAME/KEY: misc_feature
LOCATION: (51334)..(51334)
OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
LOCATION: (51602)..(51602)
OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
LOCATION: (51786)..(51786)
OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
LOCATION: (51805)..(51805)
OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
LOCATION: (55369)..(55369)
OTHER INFORMATION: n equals a, t, g or c
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NAME/KEY: misc_feature
LOCATION: (65309)..(65309)
OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
LOCATION: (65313)..(65313)
OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
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LOCATION: (122167)..(122167)
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LOCATION: (145058)..(145058)
OTHER INFORMATION: n equals a, t, g or c
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NAME/KEY: misc_feature
LOCATION: (145171)..(145171)
OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
LOCATION: (145942)..(145942)
OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
LOCATION: (147197)..(147197)
OTHER INFORMATION: n equals a, t, g or c
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NAME/KEY: misc_feature
LOCATION: (150841)..(150841)
OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature

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1 OTHER INFORMATION: n equals a, t, g or c
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4 LOCATION: (36513)..(36543)
5 OTHER INFORMATION: n equals a, t, g or c
6 FEATURE:
7 NAME/KEY: misc_feature
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10 FEATURE:
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12 LOCATION: (36536)..(36536)
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47 NAME/KEY: misc_feature
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49 OTHER INFORMATION: n equals a, t, g or c
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51 NAME/KEY: misc_feature
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55 NAME/KEY: misc_feature
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65 OTHER INFORMATION: n equals a, t, g or c
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72 LOCATION: (80024)..(80024)
73 OTHER INFORMATION: n equals a, t, g or c

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OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
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OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
LOCATION: (119750)..(119750)
OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
LOCATION: (119924)..(119924)
OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
LOCATION: (120038)..(120038)
OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
LOCATION: (121344)..(121344)
OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
LOCATION: (122167)..(122167)
OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
LOCATION: (122336)..(122336)
OTHER INFORMATION: n equals a, t, g or c
FEATURE:
NAME/KEY: misc_feature
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OTHER INFORMATION: n equals a, t, g or c
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LOCATION: (142750)..(142750)
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NAME/KEY: misc_feature
LOCATION: (145058)..(145058)
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NAME/KEY: misc_feature
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OTHER INFORMATION: n equals a, t, g or c
FEATURE:

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NAME/KEY: misc_feature
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OTHER INFORMATION: n equals a, t, g or c
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NAME/KEY: misc_feature
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LOCATION: (152530)..(152530)

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Query Match 7.8%; Score 73.6; DB 15; Length 1830121;
 Best Local Similarity 54.9%; Pred. No. 0.0034;
 Matches 167; Conservative 0; Mismatches 134; Indels 3; Gaps 1;

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QY 30 TAATGACCTTTAAATTAATGATGAGATTCAGTAGAATCTATTATTAATCAACGCTTC 89
DB 1646128 TTATTAAGCTGACCAATATATAGATGAAGCATTCATCATATTAATCAGATTATGA 1646187
QY 90 TGATTTGAGTGCATATATGTCATGATATCCAGTAGAGGATTTAAGCAATTCCT 149
DB 1646188 AATCTAGAAATTAATGCTTTCATATGCTTCAATGCTTCAACGATTTGACTTTGCTCA---TTT 1646244
QY 150 AACGAAATATTCAGTTGTAGATTAATGAATTAATAAATCTTGCTTAATGAAGAAATATTTGG 209
DB 1646245 AGAAGAAATATCTAAATTAATGAATTAAGATTAATAAATTAATCACTAATTAATTAATGAG 1646304
QY 210 TTATGACATCAAGTTTGACAAACCGGGAATTTCTTAAGGGAATATATTTTGAAT 269
DB 1646305 GTTCATTAATCTTTGATATATAGCCCTGCTGTTTTCAGGTAATTAATTTTCAAGAAAT 1646364
QY 270 GGAATGCTGATGATATTTTCATATCCAGTAGATTTGATTAAGCAATTCGTTTATGAGGA 329
DB 1646365 GGATGCTGATGATATTAATCAATCAATCGATGATTTGAAGAAATAGTTAATCTATCTGAGAA 1646424
QY 330 AAAT 333
DB 1646425 AAAT 1646428

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RESULT 9
 US-09-816-028A-30
 Sequence 30, Application US/09816028A
 Patent No. US20020042369A1
 GENERAL INFORMATION:
 APPLICANT: Gilbert, Michel
 APPLICANT: Makarchuk, Warren W.
 TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
 TITLE OF INVENTION: Gangliosides and Ganglioside Mimics
 FILE REFERENCE: 019635-000111US
 CURRENT APPLICATION NUMBER: US/09/816,028A
 PRIOR FILING DATE: 2001-03-21
 PRIOR APPLICATION NUMBER: US 60/118,213
 PRIOR FILING DATE: 1999-02-01
 PRIOR APPLICATION NUMBER: US 09/495,406
 NUMBER OF SEQ ID NOS: 49
 SOFTWARE: Patent Ver. 2.1
 SEQ ID NO 30
 TYPE: DNA
 ORGANISM: Campylobacter jejuni
 FEATURE:
 NAME/KEY: CDS

LOCATION: (1)..(891)
OTHER INFORMATION: beta-1,3 galactosyl transferase from C. jejuni O:10
US-09-816-028A-30

Query Match 7.8%; Score 73.2; DB 9; Length 891;
Best Local Similarity 49.1%; Pred. No. 0.00028;
Matches 223; Conservative 0; Mismatches 228; Indels 3; Gaps 1;

QY 1 ATGAATTAATAGTATCATATGTCGTATATATAGACCTTTAAATATGAGATTC 60
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QY 121 CCAAGTAGAGTATTAAGCAATCTTACAGATATTCAGTTAGATATAGATA 180
DB 118 TGTGAAAGTGCACAAAGTATGATATGTTAAAGATATGCCAAAAAGTATGATA 177
QY 181 AAATCTTGCTTATATAGAGAAATATGTTAGCATCAAGTTGAACAAAGCGTAAA 240
DB 178 AAATCATACATATAGAGAAATTTAAACCTTTAGAGCTGATATGAGGTGATA 237
QY 241 ATTTCTAAGGAGAAATATATTTTGAATGAGTCTGATGATATTCATATCCAGTAA 300
DB 238 GTAGCAAACTCTCTTATATATATGTTTATGATCTGATGATTTTGAAGCTTAATGCT 297
QY 301 TTGATATAGCAAAATTCCTTTATGAGAGAAATTCATGATTTCTCAGCACTTATA 360
DB 298 TGTGAAAGTATGATGAAATTTTAAACATGAAATGATTTATATTTTATGCA 357
QY 361 GAATGATAGCCAAAGAAATTTGATATATTAACAGAGAAAGTATTAATATATC 420
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QY 421 TTAATATGATATAGCAAGATGTTATGATA 454
DB 418 GTAAAAAAGATTTTAAAGAACTATTAATA 451

RESULT 10
US-10-303-161-30
Sequence 30, Application US/10303161
Publication No. US20030148459A1
GENERAL INFORMATION:
APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
TITLE OF INVENTION: National Research Council of Canada
TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
FILE REFERENCE: 019633-000111US
CURRENT APPLICATION NUMBER: US/10/303,161
CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 30
LENGTH: 891
TYPE: DNA
ORGANISM: Campylobacter jejuni
FEATURE:
NAME/KEY: CDS
LOCATION: (1)..(891)
OTHER INFORMATION: beta-1,3 galactosyl transferase from C. jejuni O:10
US-10-303-161-30

Query Match 7.8%; Score 73.2; DB 14; Length 891;

Best Local Similarity 49.1%; Pred. No. 0.00028;
Matches 223; Conservative 0; Mismatches 228; Indels 3; Gaps 1;

QY 1 ATGAATTAATAGTATCATATGTCGTATATATAGACCTTTAAATATGAGATTC 60
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DB 178 AAATCATACATATAGAGAAATTTAAACCTTTAGAGCTGATATGAGGTGATA 237
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DB 358 TTTGATATGAAATATCAATTAATGAAAGAGTGAATTTTCAAGAAAAATGTTAT 417
QY 421 TTAATATGATATAGCAAGATGTTATGATA 454
DB 418 GTAAAAAAGATTTTAAAGAACTATTAATA 451

RESULT 11
US-10-303-118-30
Sequence 30, Application US/10303118
Publication No. US20030157655A1
GENERAL INFORMATION:
APPLICANT: Gilbert, Michel
APPLICANT: Wakarchuk, Warren W.
TITLE OF INVENTION: National Research Council of Canada
TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
FILE REFERENCE: 019633-000111US
CURRENT APPLICATION NUMBER: US/10/303,118
CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 30
LENGTH: 891
TYPE: DNA
ORGANISM: Campylobacter jejuni
FEATURE:
NAME/KEY: CDS
LOCATION: (1)..(891)
OTHER INFORMATION: beta-1,3 galactosyl transferase from C. jejuni O:10
US-10-303-118-30

Query Match 7.8%; Score 73.2; DB 14; Length 891;
Best Local Similarity 49.1%; Pred. No. 0.00028;
Matches 223; Conservative 0; Mismatches 228; Indels 3; Gaps 1;

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238 GTAGCAAACTCTCTTATATATGTTTATGATCCGATGATATTTAGAACTTATCT 297
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RESULT 14
US-10-303-162-30
Sequence 30, Application US/10303162
Publication No. US20030157658A1
GENERAL INFORMATION:
APPLICANT: Gilbert, Michel
APPLICANT: Makarchuk, Warren W.
APPLICANT: National Research Council of Canada
TITLE OF INVENTION: Campylobacter Glycosyltransferases for Biosynthesis of
TITLE OF INVENTION: Gangliosides and Ganglioside Mimics
FILE REFERENCE: 019633-000111US
CURRENT APPLICATION NUMBER: US/10/303,162
CURRENT FILING DATE: 2002-11-21
PRIOR APPLICATION NUMBER: US/09/816,028
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: US 60/118,213
PRIOR FILING DATE: 1999-02-01
PRIOR APPLICATION NUMBER: US 09/495,406
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 49
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 30
LENGTH: 891
TYPE: DNA
ORGANISM: Campylobacter jejuni
FEATURE:
NAME/KEY: CDS
LOCATION: (1)..(891)
OTHER INFORMATION: beta-1,3 galactosyl transferase from C. jejuni O:10
US-10-303-162-30
Query Match 7.8%; Score 73.2; DB 14; Length 891;
Best Local Similarity 49.1%; Pred. No. 0.00028; Indels 3; Gaps 1;
Matches 223; Conservative 0; Mismatches 228;
QY 1 ATGATATATGATCATATATGTCGATATATATGAGCCCTTAAATATATGAGAGATCA 60
1 ATGTTTAAATTTCAATCATCTTCCCACTTATATATGAGAACATATATAGCAAGGCA 60
Db
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61 ATGAAAAAGTTGTAACAATCAAGATTTTAAAAATATGA--AAATATGATGATGAT 117
Db
QY 121 CCAAGTAGAGGTGATTTAAAGCAATTCCTTAACAGAAATTCAGTTGATATAGATA 180
118 TGTGAGAGTACAAAAGATATGATATATGAGAAATATGCAAAAAGATGATAGATA 177
QY 181 AAAATCTGCTTAATGAGAGAAATTTGCTTTAGCATCAAGTTGAAACAAAGCGTGA 240
178 AAAATCATATCATATGAGAGAAATTTTAAACTTTTAAAGAGCTAGATATGAGAGGTGAAA 237
QY 241 ATTCTAAGAGAGATATATTTTATGAGATGCTGATGATATTTATATATCCAAAGTGA 300

Db 238 GTAGCAAACTCTCTTATATATGTTTATGATCCGATGATATTTAGAACTTATGCT 297
QY 301 TTGATTAAGCAAAATGCTTTTATGAGAGAAATTCATGATTTCTCAAGCTTAA 360
298 TGTGAGAGATGATGAGAAATTTTAAACATGAGAAATGATTTTATTTTATGCA 357
QY 361 GAATGATGACCAAAAGAAATTTGATATATTAACAAGAGATATATTAATATAC 420
358 TTGATTAAGCAAAATTAACAATTAAGAAAGATTTTCAAGAAATGTTAT 417
QY 421 TTACTAATGATATAGGAGATGTTATGATA 454
418 GTAAAAAAGATTTTAAAAAGACTATTTAAAA 451
Db
RESULT 15
US-10-282-122A-17153
Sequence 17153, Application US/10282122A
Publication No. US20040029129A1
GENERAL INFORMATION:
APPLICANT: Wang, Liangsu
APPLICANT: Zamudio, Carlos
APPLICANT: Malone, Cheryl
APPLICANT: Haselbeck, Robert
APPLICANT: Ohlsen, Kari
APPLICANT: Zykkind, Judith
APPLICANT: Wall, Daniel
APPLICANT: Trawick, John
APPLICANT: Carr, Grant
APPLICANT: Yamamoto, Robert
APPLICANT: Forsyth, R.
APPLICANT: Xu, H.
TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
FILE REFERENCE: EUTTRA-034A US/10/282,122A
CURRENT APPLICATION NUMBER: 2003-02-20
CURRENT FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: 60/206,848
PRIOR FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: 60/230,335
PRIOR FILING DATE: 2000-09-06
PRIOR APPLICATION NUMBER: 60/230,347
PRIOR FILING DATE: 2000-09-09
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR FILING DATE: 2000-10-23
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR FILING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR FILING DATE: 2000-12-22
PRIOR APPLICATION NUMBER: 60/267,636
PRIOR FILING DATE: 2001-02-09
PRIOR APPLICATION NUMBER: 60/269,308
PRIOR FILING DATE: 2001-02-16
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 78614
SOFTWARE: PatentIn version 3.1
SEQ ID NO 17153
LENGTH: 747
TYPE: DNA
ORGANISM: Clostridium difficile
US-10-282-122A-17153
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Best Local Similarity 51.3%; Pred. No. 0.00053;
Matches 192; Conservative 0; Mismatches 179; Indels 3; Gaps 1;
QY 32 ATGAGCCTTAATATATGATGAGAGATTCAGTATGATATATTAATCAAGCGTATCG 91
38 ATATATCGAAAAATTTATGAGAACCAATTAATATGATTAATCAAGCGTATCG 97

QY 92 ATTTGAGTTCATATGTCATTTGATATCCAGTAGAGTGATTTAAGCAATTCCTTA 151
 Db 98 AATGGGAATGTTAAATTT--ATTGATGATTTGCTCAACGATATAGTCTTAATAGTCA 154
 QY 152 CAGATATTCAGTTGTAGATATAGATATTAATTTCTTAATGAGAAATATTTGTT 211
 Db 155 AATCTTATATGCAACAGATAGTAGATATTAATATGATTAAGACTGAGACTAATAGGCTG 214
 QY 212 TAGCATCAAGTTTGAACAAGGGTGAAATTTCTAAGGAGATATATTTTAGAATCG 271
 Db 215 TCTTAATGCTAGAAATTTAGCACTAAGTAGGCAACGACAAATTTATAGCTTTTATG 274
 QY 272 ATGCTGATGATATTTCAATCCAGTAGATTTGATAGCAATTCGTTTATGAGAGAA 331
 Db 275 ATAGTATGACCAATGGAATAGTAGTAAGTGAAGAAACAGTAATTTATGTTAGAA 334
 QY 332 ATTCATGATTTCTCAGCACTCTATAGAAATTTGATAGACCAAAAGAAATTTAGAT 391
 Db 335 ATGACTATGTAAATTTCAATTTACTTCAATGAACTGATGAAATGATTAATAATTA 394
 QY 392 ATAAACAACGAGAA 405
 Db 395 ACAAGTAAATAAA 408

Search completed: March 28, 2004, 08:50:48
 Job time : 438 secs